

5891432

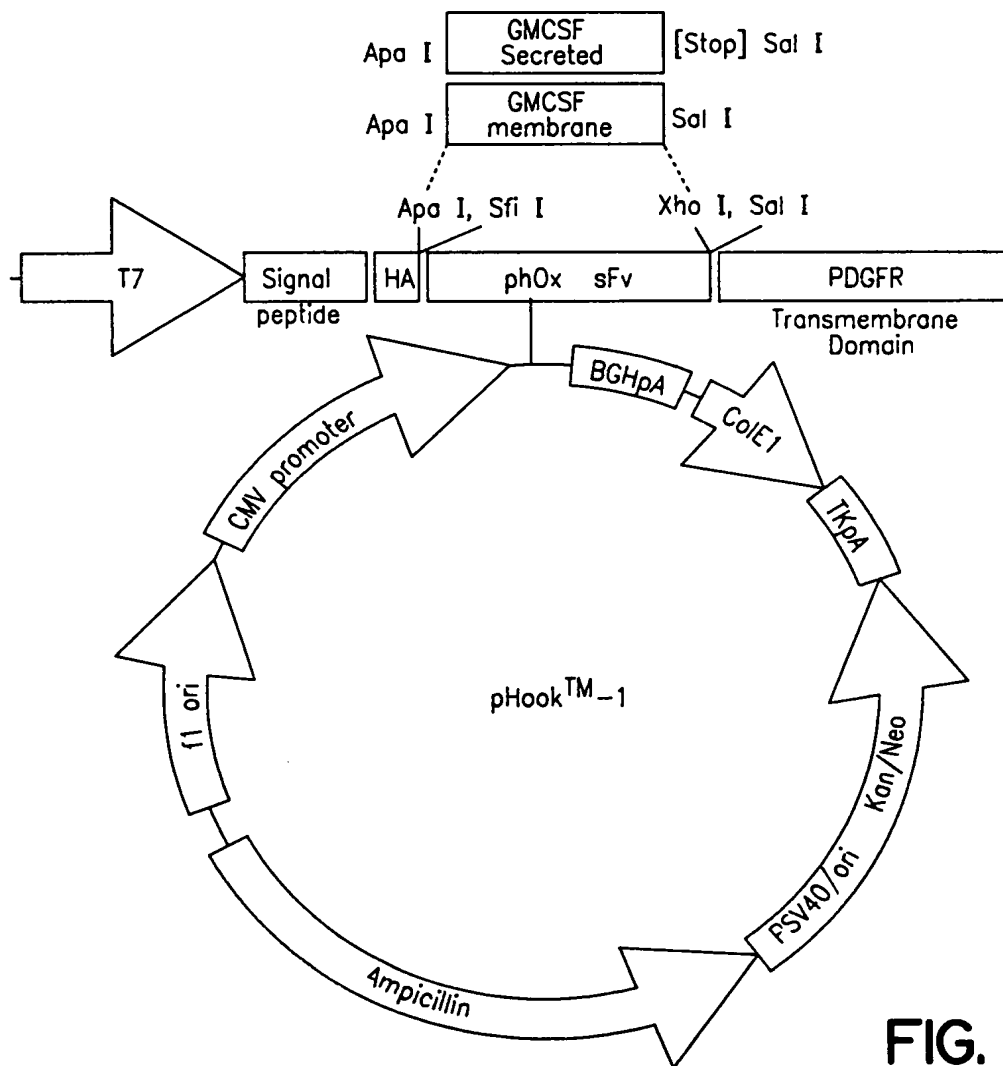


FIG. 1A

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
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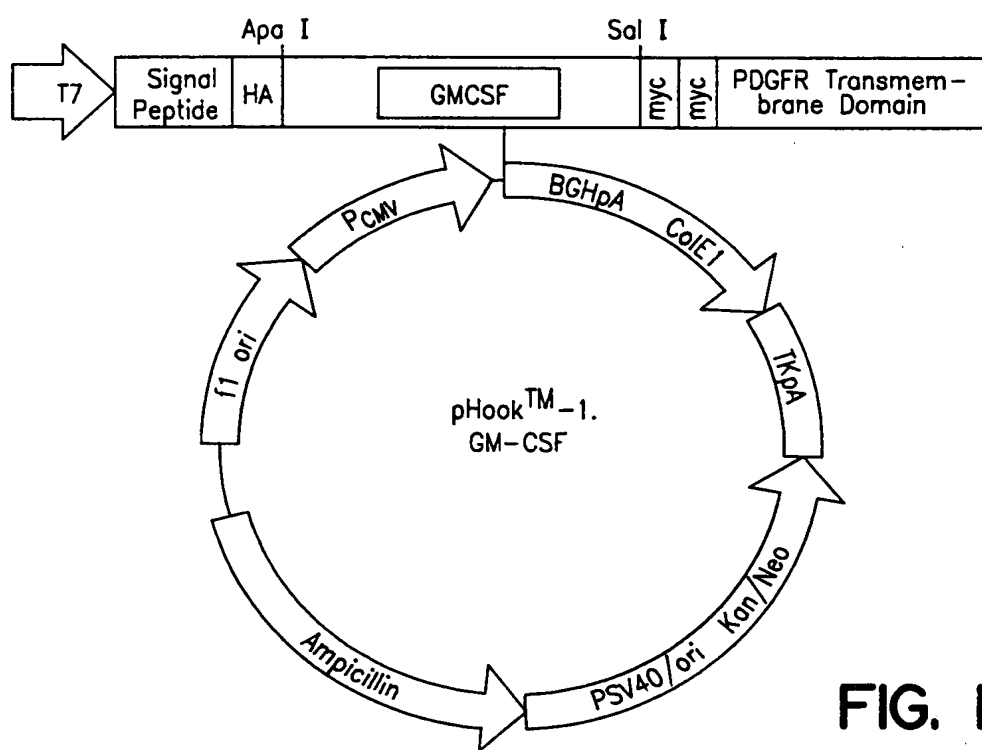


FIG. 1B

APPROVED	O.G. FIG.	
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      360      370      380      390      400
      *  *      *  *      *  *      *  *      *  *
CTGCCCCCCA ACTCCGGAAA CGGACTGTGA AACACAAGTT ACCACCTATG
  C  P  P    T  P  E    T  D  C  E    T  Q  V    T  T  Y

      410      420      430      440      450
      *  *      *  *      *  *      *  *      *  *
CGGATTTTCAT AGACAGCCTT AAAACCTTTC TGA CTGATAT CCCCTTTGAA
  A  D  F  I    D  S  L    K  T  F    L  T  D  I    P  F  E

      460      470      480      490      500
      *  *      *  *      *  *      *  *      *  *
TGCAAAAAAC CAGGCCAAAA AGTCGACGAA CAAAAACTCA TCTCAGAAGA
  C  K  K    P  G  Q  K    V  D  E    Q  K  L    I  S  E  E
— mouse GMCSF —>

      510      520      530      540      550
      *  *      *  *      *  *      *  *      *  *
GGATCTGAAT GCTGTGGGCC AGGACACGCA GGAGGTCATC GTGGTGCCAC
  D  L  N    A  V  G    Q  D  T  Q    E  V  I    V  V  P

      560      570      580      590      600
      *  *      *  *      *  *      *  *      *  *
ACTCCTTGCC CTTTAAGGTG GTGGTGATCT CAGCCATCCT GGCCCTGGTG
  H  S  L  P    F  K  V    V  V  I    S  A  I  L    A  L  V
                    <— PDGFRβ transmembrane domain...—

      610      620      630      640      650
      *  *      *  *      *  *      *  *      *  *
GTGCTCACCA TCATCTCCCT TATCATCCTC ATCATGCTTT GGCAGAAGAA
  V  L  T    I  I  S  L    I  I  L    I  M  L    W  Q  K  K
transmembrane domain—

      660      670      680      690
      *  *      *  *      *  *      *  *
GCCACGTTAG GCGGCCGCTC GAGATCAGCC TCGACTGTGC CTTCTAG
  P  R  *    A  A  A    R  D  Q  P    R  L  C    L  L
—>

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FIG. 2B

APPROVED	O.G. FIG.	
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1 2 3 4 5 6 7 8 9 10 11 12 13 14 15

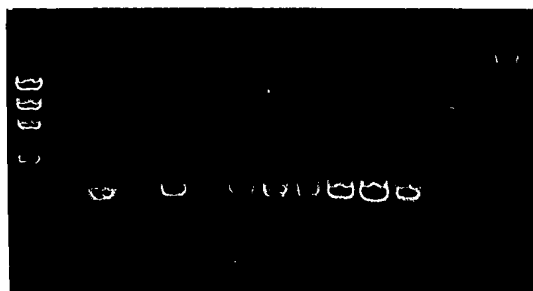


FIG. 3

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
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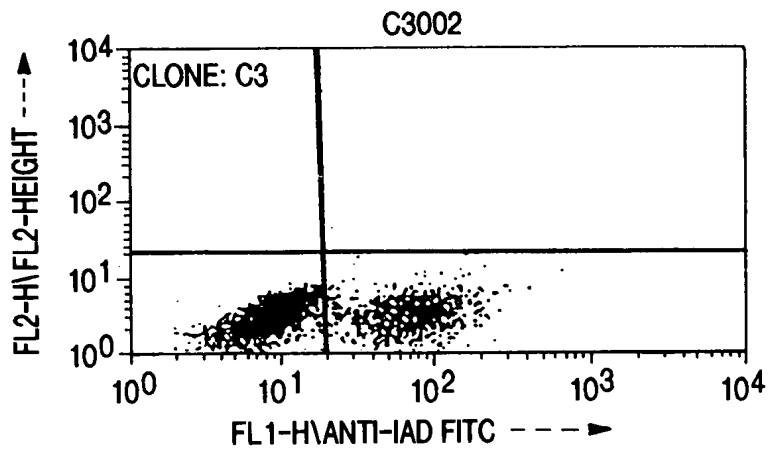


FIG. 4A

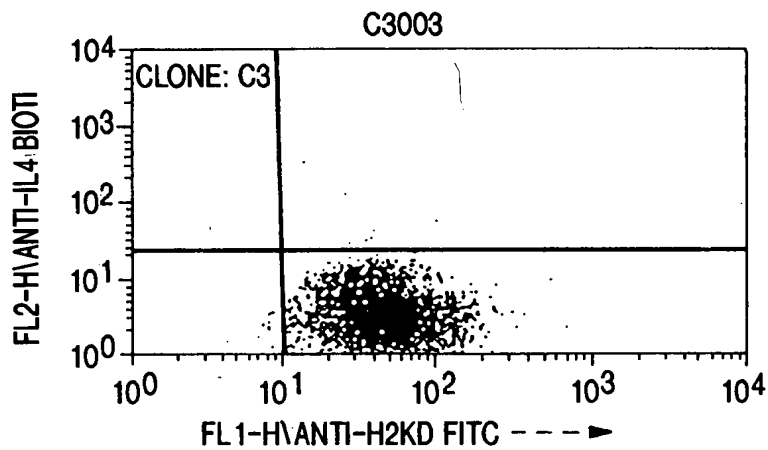


FIG. 4B

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
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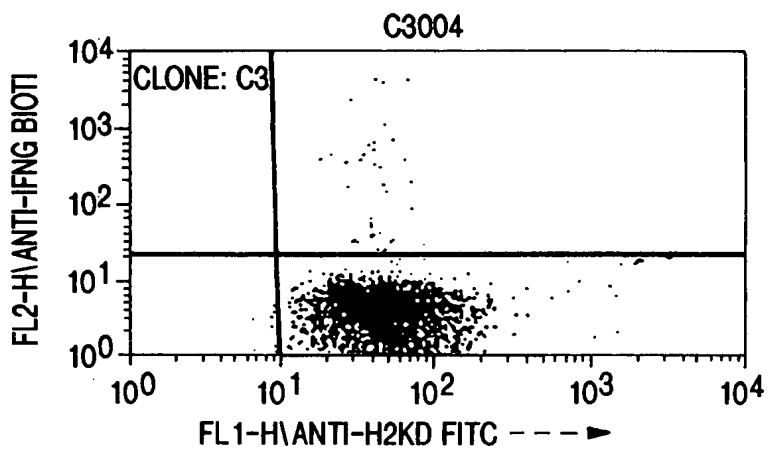


FIG. 4C

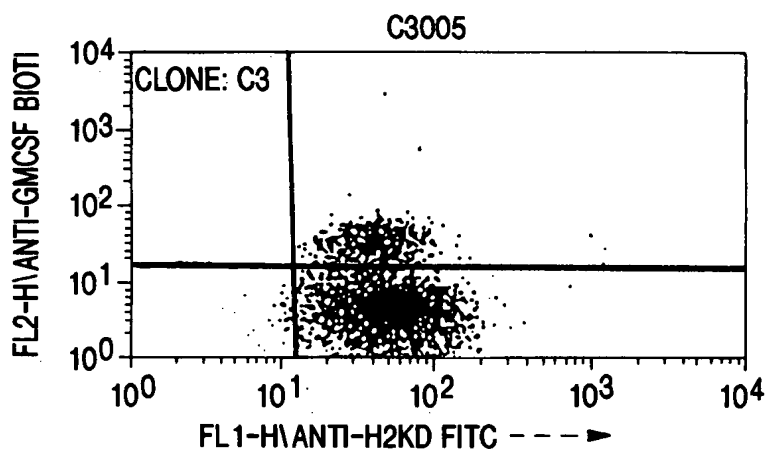


FIG. 4D